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Sequence Listing was accepted.

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Reviewer: Keisha Douglas

Timestamp: Thu Sep 27 13:33:56 EDT 2007

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Application No: 10579614 Version No: 1.1

Input Set:

Output Set:

Started: 2007-09-27 13:33:01.977
Finished: 2007-09-27 13:33:17.030
Elapsed: 0 hr(s) 0 min(s) 15 sec(s) 53 ms
Total Warnings: 62
Total Errors: 0
No. of SeqIDs Defined: 97
Actual SeqID Count: 97

Error code	Error Description
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W 402	Undefined organism found in <213> in SEQ ID (3)
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W 402	Undefined organism found in <213> in SEQ ID (23)
W 402	Undefined organism found in <213> in SEQ ID (25)
W 402	Undefined organism found in <213> in SEQ ID (26)
W 402	Undefined organism found in <213> in SEQ ID (27)

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Error code	Error Description
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SEQUENCE LISTING

<110> De Jong, Jan C.
Besteboer, Theodorus M.
Simon, James H.M.
Fouchier, Ronaldus A.M.
Osterhaus, Albertus D.M.E.

<120> Novel atypical pneumonia-causing virus

<130> P67119US00

<140> 10579614
<141> 2006-05-17

<150> EP 03078613.1
<151> 2003-11-18

<150> EP 03078772.5
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<150> PCT/NL2004/000805
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10 15 20 25
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Val Ala Val Arg Ala Tyr Ser Glu Ala Ala Ala Gln Gly Phe Gln Ala
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tgc cgc ttt gtt gct ttt ggc tta cag gat tgt gta acc ggt att aat 435
Cys Arg Phe Val Ala Phe Gly Leu Gln Asp Cys Val Thr Gly Ile Asn
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Asp Asp Asp Tyr Val Ile Ala Leu Thr Gly Thr Asn Gln Leu Cys Ala
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Phe Gly Ser Pro Phe Ile Thr Asn Gly Ile Ser Leu Leu Asp Ile Ile				
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Tyr Ala Gly Leu Val Val Lys His Ile Thr Asn Ile Thr Gly Val Ser				
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365	370	375		
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Ile Asp Ile Ser Ala Gly Met Phe Gly Leu Tyr Asp Asp Ile Leu Thr				
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Asn Asn Lys Pro Trp Phe Val Arg Lys Ala Ser Gly Leu Phe Asp Ala				
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430	435	440	
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445	450	455	
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Gln Pro Val Tyr Arg Thr Phe Thr Gln Ala Ile Cys Ala Ala Phe Asp			
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Asp Ala Ile Ile Ala Val Glu Leu Leu Ser Asp Phe Lys Thr Ala			
620	625	630	

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Val Phe Val Tyr Thr Cys Val Val Asp Gly Cys Ser Val Ile Val Arg			
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860	865	870	
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Tyr Val Val Asp Asp Ile Tyr Tyr Pro Ala Ser Cys Asn Gly Val Leu			
875	880	885	
cca gtt gct ttt aca aaa ttg gca ggt ggt aaa ata tct ttt tct gat			2979
Pro Val Ala Phe Thr Lys Leu Ala Gly Gly Lys Ile Ser Phe Ser Asp			
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Asp Val Ile Val His Asp Val Glu Pro Thr His Lys Val Lys Leu Ile			
910	915	920	
ttt gag ttt gaa gat gat gtt acc agt ctt tgt aag aag agt ttt			3075
Phe Glu Phe Glu Asp Asp Val Val Thr Ser Leu Cys Lys Lys Ser Phe			
925	930	935	
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